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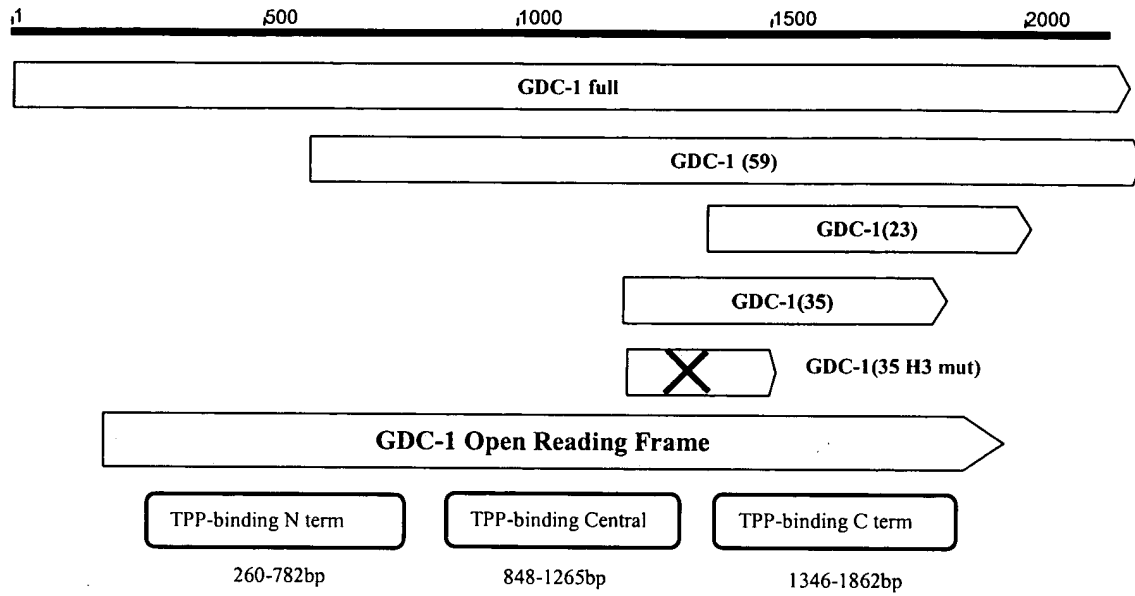


FIG. 1

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      *      20      *      40      *      60      *
GDC-1(full) : MASINIRVQNLEQPMQVAEYLFRRLEIGIRSIHGLPGDYNPLALDYLPSGCLRUVGSGVNELNAAYAADGYARVK : 75
GDC-1_(23) : ----- : -
GDC-1_(35) : ----- : -
GDC-1_(59) : ----- : -

      *
      80      *      100      *      120      *      140      *
GDC-1(full) : QMGALITTFVGVGELSAINGVAGAFSEHVPPVHIVGCPSTASQRNGMLLHHTLGNQDFNIFANMSAQISCEVAKLT : 150
GDC-1_(23) : ----- : -
GDC-1_(35) : ----- : -
GDC-1_(59) : -----RNGMLLHHTLGNQDFNIFANMSAQISCEVAKLT : 33

      160      *      180      *      200      *      220
GDC-1(full) : NPAETATQIDHALRVCFIRSRPVYIHLPTDMVOAKVEGARLKEPIDLSEPPNDPEKEAYVVDVVLKYLRAAKNPV : 225
GDC-1_(23) : ----- : -
GDC-1_(35) : ----- : -
GDC-1_(59) : NPAETATQIDHALRVCFIRSRPVYIHLPTDMVOAKVEGARLKEPIDLSEPPNDPEKEAYVVDVVLKYLRAAKNPV : 108

      *      240      *      260      *      280      *      300
GDC-1(full) : ILVDACAIRHRVLDDEVHDLIEKTINLPVFVTPMGKGAVNEEHPTYGGVYAGDGSHPPOVKDMVESSDLILITGALK : 300
GDC-1_(23) : ----- : -
GDC-1_(35) : ----- : -
GDC-1_(59) : ILVDACAIRHRVLDDEVHDLIEKTINLPVFVTPMGKGAVNEEHPTYGGVYAGDGSHPPOVKDMVESSDLILITGALK : 183

      *      320      *      340      *      360      *
GDC-1(full) : SDFTAGFSYRTSQLNTIDLHSDHCIVKYSTYPGVQMRGVLRQVIKQLDASEINAQPAPVVENEVAKNRDMSFVI : 375
GDC-1_(23) : ----- : -
GDC-1_(35) : ----- : -
GDC-1_(59) : SDFTAGFSYRTSQLNTIDLHSDHCIVKYSTYPGVQMRGVLRQVIKQLDASEINAQPAPVVENEVAKNRDMSFVI : 258
                      typgvqmrqvlrqvikqldaseinaqpapvvenevaknrdsnspvi

      380      *      400      *      420      *      440      *
GDC-1(full) : TQAFFWPRVGEFLKKNDIVITETGTANFGIWDTKFPSCVTALSQVLWGSIGWSVGACQGAFLAAADDMSDRRTIL : 450
GDC-1_(23) : ---FFWPRVGEFLKKNDIVITETGTANFGIWDTKFPSCVTALSQVLWGSIGWSVGACQGAFLAAADDMSDRRTIL : 72
GDC-1_(35) : TQAFFWPRVGEFLKKNDIVITETGTANFGIWDTKFPSCVTALSQVLWGSIGWSVGACQGAFLAAADDMSDRRTIL : 120
GDC-1_(59) : TQAFFWPRVGEFLKKNDIVITETGTANFGIWDTKFPSCVTALSQVLWGSIGWSVGACQGAFLAAADDMSDRRTIL : 333
                      tqaffwprvgeflkkndivitetgtanfgiwdtkfpscvtalsqvlwgsigwsvgacqgaflaaaddmsdrtil

      460      *      480      *      500      *      520
GDC-1(full) : FVGDGSFQLTAQELSTHIRLKLKPIIFVICNDGFTIERFIHGHEAEYNDIAMWDFKALVDVFGGSKTAKKFAVKT : 525
GDC-1_(23) : FVGDGSFQLTAQELSTHIRLKLKPIIFVICNDGFTIERFIHGHEAEYNDIAMWDFKALVDVFGGSKTAKKFAVKT : 147
GDC-1_(35) : FVGDGSFQLTAQELSTHIRLKLKPIIFVICNDGFTIERFIHGHEAEYNDIAMWDFKALVDVFGGSKTAKKFAVKT : 195
GDC-1_(59) : FVGDGSFQLTAQELSTHIRLKLKPIIFVICNDGFTIERFIHGHEAEYNDIAMWDFKALVDVFGGSKTAKKFAVKT : 408
                      FVGDGSFQLTAQELSTHIRLKLKPIIFVICNDGFTIERFIHGHEAEYNDIAMWDFKALVDVFGGSKTAKKFAVKT

      *      540      *      560      *
GDC-1(full) : KDELDSSLTDPTFMAAECLQFVELYMPKEDAPRALINTAEASARNNAKTE : 575
GDC-1_(23) : KDELDSSLTDPTFMAAECLQFVELYMPKEDAPRALINTAEASARNNAKTE : 197
GDC-1_(35) : KDELDSSLTDPTFMAAECLQFVELYMPKEDAPRALINTAEASARNNAKTE : 226
GDC-1_(59) : KDELDSSLTDPTFMAAECLQFVELYMPKEDAPRALINTAEASARNNAKTE : 458
                      KDELDSSLTDPTFMAAECLQFVELYMPKEDAPRALINTAEASARNNA*te

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FIG. 2

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      *      20      *      40      *      60      *
GDC-1(full) : MASINIRVONLEQPMQAEYLRRLHEVIGIRSIHCLPGDYMLALDYLPSCG-LEMYGCVMLNAAYAADGYARV : 74
A_oryzae-PDC : -----MSISTSSGDFVRUAFVQYIVFVVSINAPAGVYMLVALDYLPKED-LEMYGCVMLNAGYAADGYARI : 66
E._nidulans_PDC : --MADIATRDVROPIDIAEYLRRLHEVIGIRSVHGVPGDYMLAALDYLPKED-LEMYGCVMLNAGYAADGYARV : 72
S.cer_PDC1 : -----MSEITLGGKYLPERLKQVNNVNTVFLPGDFMLSLDDKIYEVECHRMAGNAGNELNARYAADGYARI : 64
S.cer_PDC5 : -----MSEITLGGKYLPERLSQVNCNNTVFLPGDFMLSLDDKIYEVECHRMAGNAGNELNARYAADGYARI : 64
              yLf r1 6 36 g pgD5N* L1 6 6 W Gn NELNA YAADGYAR6

      80      *      100      *      120      *      140      *
GDC-1(full) : KMGALITTFGVGELSALMGVAGAFSEHWVPHVHVGCPSTASQRNGHLLHHTLGNMGDFNIFRAMSAQISCEVAKL : 149
A_oryzae-PDC : NMGSAVLTTFGVGELSALMALAGAFSEFVPIVHIVGQEHTEKQKQDGHLLHHTLGNMGDFNIVFTRMSADISCLGCL : 141
E._nidulans_PDC : NGLAALVTTFGVGELSALMALAGAFSEFVPIIHHVVGQPHSRQKDCLLHHTLGNMGDYNVFSNNKGLISVITIANL : 147
S.cer_PDC1 : KMGSCITTFGVGELSALMGVAGAFSEHWVPHVHVGCPSTASQRNGHLLHHTLGNMGDFNIVFTRMSADISCEVAKL : 139
S.cer_PDC5 : KMGSCITTFGVGELSALMGVAGAFSEHWVPHVHVGCPSTASQRNGHLLHHTLGNMGDFNIVFTRMSADISCEVAKL : 139
              g6 66TTFGVGELSA6N 6AG 5 E V 66H6VG P *5Q 6LLHHTLGNMG5 6F Msa IS t a 6

      160      *      180      *      200      *      220
GDC-1(full) : EMPAEIATGLIHAFVCFTRSRPVYIMLFDIMVQAKVGGARLKEPIDLSEFPMDPEKAYVVDVWVLYKYLRA--AF : 222
A_oryzae-PDC : NSTHEVATLIDNAIFEGMTSRPVYISLPDMVTKIKIEGERIDTFLDLSEFPMDPEKEDYVVDVWVLYKYLHA--AF : 214
E._nidulans_PDC : NPTYDAATLIDNAIFEGMTSRPVYIALFDMITKKIEGERIKTPIDLSEFPMDPEKEDYVVDARAQVFAFSAR : 222
S.cer_PDC1 : IDICTAPAEIDRCITITTYTORPVYLGPAHLVGLNVPKAKLQTPIDHSILKPHDAESEKEVITDTILVLAKD--AF : 212
S.cer_PDC5 : IDIRNAPAEIDRCITITTYTORPVYLGPAHLVGLNVPKAKLQTPIDLSILKPHDAEAEVVRTVVEFIKD--AF : 212
              ID 6R 5 RPYV6 LP 166 6 L tP6D6S1 pND E E V6 AK

      *      240      *      260      *      280      *      300
GDC-1(full) : NPVILVDACAIRHR---WDEVDHDIIEKIMLPEVFI PHGKCAVNEEHETTYGCVYACDGHPPQVDMWSESDLI : 293
A_oryzae-PDC : KPVILVDACAIRHR---WDEVHEFPWEKSGLPFVAPHGKCAVDETHKNYGCYVYACTGSMF-CVREQWESDLI : 284
E._nidulans_PDC : NPVILVDSEVFDSTGLSEEDHDIIEVSGLPFVAPHGKCAVNEHTRCYGCVYACTGSMF-CVREQWESDLI : 296
S.cer_PDC1 : NPVILADACESRHD---WKAEIKKLDLQFFPAFVPHGKCSI SEQHRYGCVYVETLSKP-EVREAVESADLI : 282
S.cer_PDC5 : NPVILADACESRHD---WKAEIKKLDLQFFVFWPHGKCAIDEQHRYGCVYVETLSRP-EVVKAVESADLI : 282
              nPVIL Dac rh v E 16 3 P 5V PMGKa6 E H YGGVY Gt S P V4 VES DLI

      *      320      *      340      *      360      *
GDC-1(full) : LTIGALKSDFNTAGFSYRTSQMLTIDLHSDHCIMVYSYTPGVQHGVLRQVVKQLDASEINAOAPV-VNEVEAK : 367
A_oryzae-PDC : LSIGALKSDFNTAGFSYRIGQLTIDHSTYVVRVYSEYFDIMHCEVLQRIWQRMGNINVGVPVSPPS-NLLEDNE : 358
E._nidulans_PDC : LSIGALKSDFNTAGFSYRIGQLTIDHTTYVVRVYSEYFDIMHCEVLQRIWQRMGNINVGVPVSPPS-NLLEDNE : 370
S.cer_PDC1 : LSVGALLSDFNTAGFSYRTSQMLTIDHSDHCIMVYSYTPGVQHGVLRQVVKQLDASEINAOAPV-VNEVEAK : 357
S.cer_PDC5 : LSIGALLSDFNTAGFSYRTSQMLTIDHSDHCIMVYSYTPGVQHGVLRQVVKQLDASEINAOAPV-VNEVEAK : 357
              L36GA6 SDFNT FSY N 6 FH3 64 5P M4 vL k66 6 p p

      380      *      400      *      420      *      440      *
GDC-1(full) : NRDNSPVITQAFVFPVGGFLKNDIVINDETGTANFGIUDKFFSCVYALSQVLWGSIGYSVGAECGAVLAAADD : 442
A_oryzae-PDC : KASTEQALPHANLMTVCGNLKQKQVVIETGTANFGIUDKFFPAGVTALISQVLWGSIGYSVGAECGAAALAAKQ : 433
E._nidulans_PDC : KNSSEQRIHANLMTVCGNLKQKQVVIETGTANFGICUNLLPELQPIQSQVLWGSIGYSVGAECGAAALAAKQ : 445
S.cer_PDC1 : AVPASTPKQEMMMNOLGNFLQGGVYIAETGSAFGINQITFPNNTYGISQVLWGSIGFTTGATLCAAFAAEBI : 432
S.cer_PDC5 : STPANTPKQEMMMNOLGNFLQGGVYIAETGSAFGINQITFPDQVYIVQVLWGSIGFTTGATLCAAFAAEBI : 432
              6 5 W 6G 5L e D6VI ETGT FGI t fP 6sQVLWGSIGS3vGA GA AA e

      460      *      480      *      500      *      520
GDC-1(full) : MSDRFVILFVGDGSEQLTAEISTHIRLKKPFIIFVIMDGYTIERFIHGCPAEVYHDIAMNDFKALVDVFGSKT : 517
A_oryzae-PDC : G--RFVILFVGDGSEQLTAEISTHIRNLMNPFIFVIMDGYTIERFIHGCPAEVYHDIQPMDFENIPVAFGRKQ : 506
E._nidulans_PDC : GN--RFVILWVGDSSEQLTAEISTHIRNLMNPFIFVIEGEGYTIERFIHGWDSEYHDIQPMDFENIPVAFGRGR : 519
S.cer_PDC1 : DPKRFVILFVGDGSEQLTAEISTHIRLKKPFIIFVIMDGYTIERFIHGCPAEVYHDIQPMDFENIPVAFGRKQ : 506
S.cer_PDC5 : DPKRFVILFVGDGSEQLTAEISTHIRLKKPFIIFVIMDGYTIERFIHGCPAEVYHDIQPMDFENIPVAFGRKQ : 506
              4R 6L56GDG5 QLT QE6STMR L P 6FV6 n G5TIE4 IHG a YN Iq WD 6 FG

      *      540      *      560      *      580
GDC-1(full) : AKKFAKQKQEDSLLTQPTNAAEGIFQVEYMPKEDAPRAEDTAEASARNNAKTE : 575
A_oryzae-PDC : YKGAVVITRDEIRELFAMEEFASAPGLQFEIDSRIGR----- : 543
E._nidulans_PDC : YKGAVVITRDEITKLFASEEFSSTPELQVSLCDITHYLSDAFQELG----- : 566
S.cer_PDC1 : YETHRVATGGEWDKLTQDKSFNDNSKIRHLEVMEVFDAPONLYEQAQLTAATNAKQ- : 563
S.cer_PDC5 : YETHRVATGGEWEKLTQDKSFNDNSKIRHLEVMEVFDAPONLYEQAQLTAATNAKQ- : 563
              y V T E L F 6

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FIG. 3

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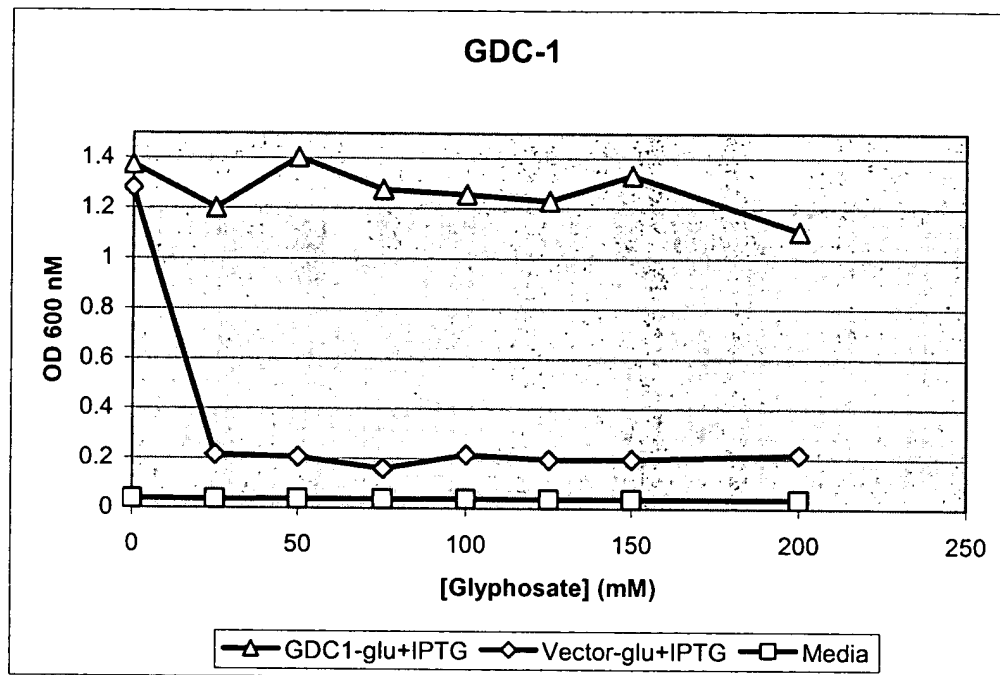
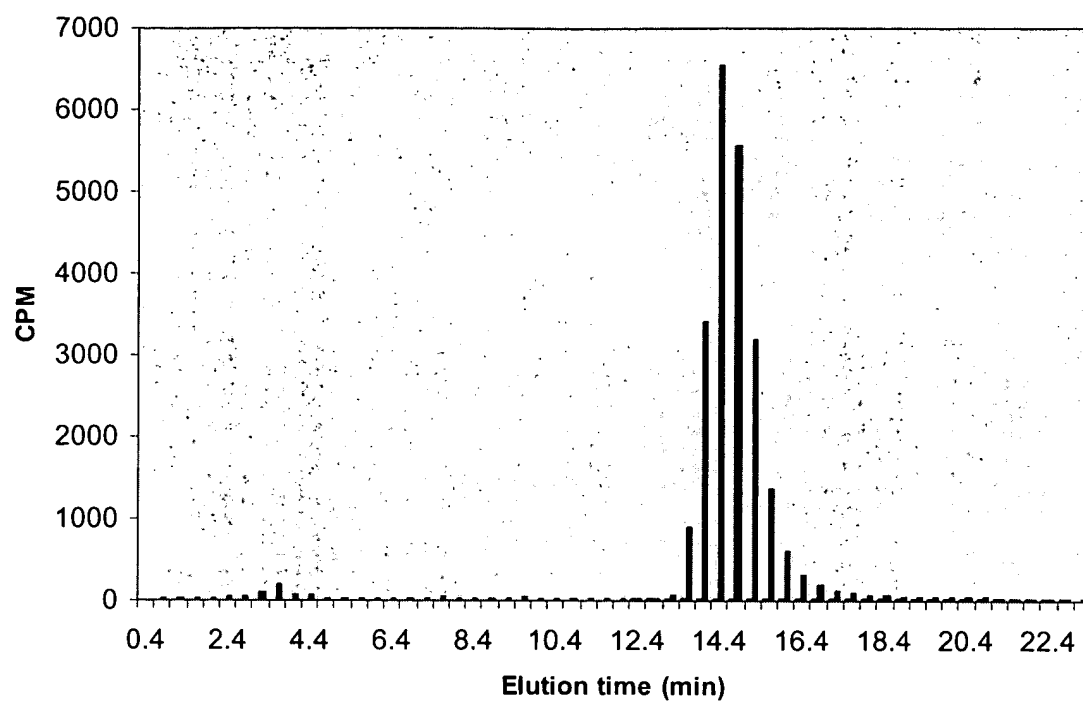


FIG. 4

A.

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B.

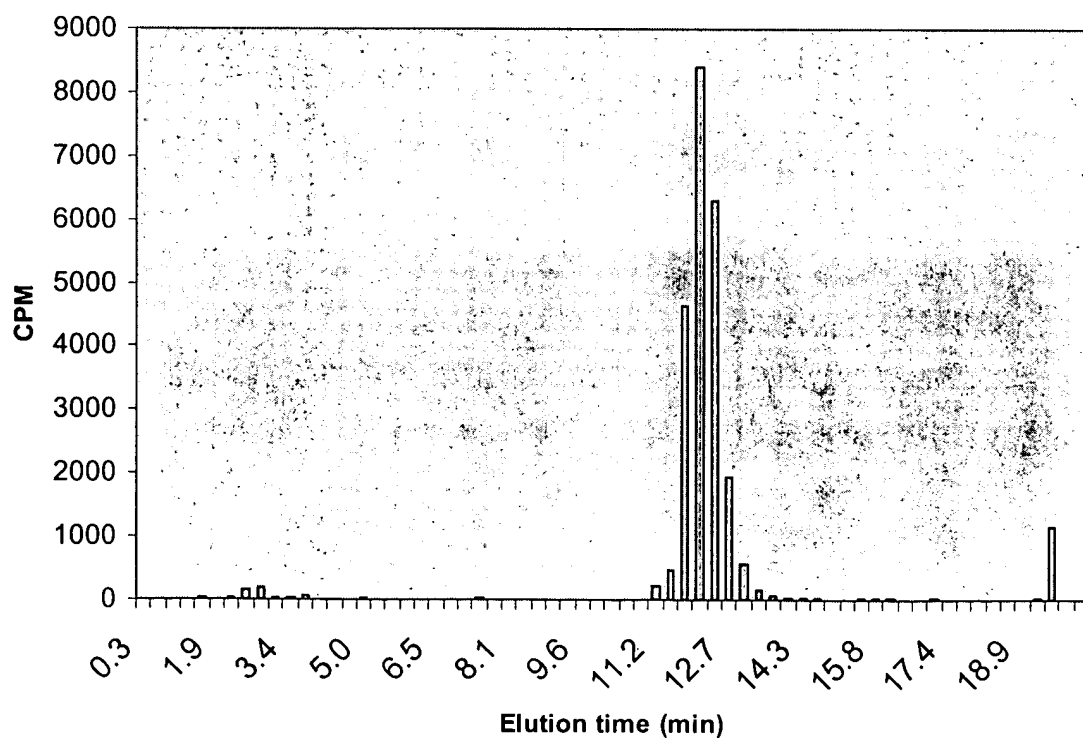


FIG. 5